

SEQUENCE LISTING

<110> Reiner, Peter B.
Connop, Bruce P.
Pollard, Michelle

<120> REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION
BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR ACTIVITY

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<141> 2002-02-08

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Glu	Glu	Thr	Ile	Ser	Ala	Met	Lys	Thr	Val	Arg	Ser	Phe	Ala	Asn	Glu
370						375					380				
Glu	Glu	Glu	Ala	Glu	Val	Tyr	Leu	Arg	Lys	Leu	Gln	Gln	Val	Tyr	Lys
385					390					395					400
Leu	Asn	Arg	Lys	Glu	Ala	Ala	Ala	Tyr	Met	Tyr	Tyr	Val	Trp	Gly	Ser
			405					410						415	
Gly	Leu	Thr	Leu	Leu	Val	Val	Gln	Val	Ser	Ile	Leu	Tyr	Tyr	Gly	Gly
			420					425					430		
His	Leu	Val	Ile	Ser	Gly	Gln	Met	Thr	Ser	Gly	Asn	Leu	Ile	Ala	Phe
	435						440					445			
Ile	Ile	Tyr	Glu	Phe	Val	Leu	Gly	Asp	Cys	Met	Glu	Ser	Val	Gly	Ser
450						455				460					
Val	Tyr	Ser	Gly	Leu	Met	Gln	Gly	Val	Gly	Ala	Ala	Glu	Lys	Val	Phe
465					470					475					480
Glu	Phe	Ile	Asp	Arg	Gln	Pro	Thr	Met	Val	His	Asp	Gly	Ser	Leu	Ala
			485					490						495	
Pro	Asp	His	Leu	Glu	Gly	Arg	Val	Asp	Phe	Glu	Asn	Val	Thr	Phe	Thr
		500						505					510		
Tyr	Arg	Thr	Arg	Pro	His	Thr	Gln	Val	Leu	Gln	Asn	Val	Ser	Phe	Ser
	515					520						525			
Leu	Ser	Pro	Gly	Lys	Val	Thr	Ala	Leu	Val	Gly	Pro	Ser	Gly	Ser	Gly
530						535				540					
Lys	Ser	Ser	Cys	Val	Asn	Ile	Leu	Glu	Asn	Phe	Tyr	Pro	Leu	Glu	Gly
545					550					555					560
Gly	Arg	Val	Leu	Leu	Asp	Gly	Lys	Pro	Ile	Ser	Ala	Tyr	Asp	His	Lys
			565					570						575	
Tyr	Leu	His	Arg	Val	Ile	Ser	Leu	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe
		580					585						590		
Ala	Arg	Ser	Ile	Thr	Asp	Asn	Ile	Ser	Tyr	Gly	Leu	Pro	Thr	Val	Pro
	595					600						605			
Phe	Glu	Met	Val	Val	Glu	Ala	Ala	Gln	Lys	Ala	Asn	Ala	His	Gly	Phe
610					615					620					
Ile	Met	Glu	Leu	Gln	Asp	Gly	Tyr	Ser	Thr	Glu	Thr	Gly	Glu	Lys	Gly
625					630					635					640
Ala	Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Val	Ala	Met	Ala	Arg	Ala
			645					650						655	
Leu	Val	Arg	Asn	Pro	Pro	Val	Leu	Ile	Leu	Asp	Glu	Ala	Thr	Ser	Ala
		660					665						670		
Leu	Asp	Ala	Glu	Ser	Glu	Tyr	Leu	Ile	Gln	Gln	Ala	Ile	His	Gly	Asn
	675					680						685			
Leu	Gln	Lys	His	Thr	Val	Leu	Ile	Ile	Ala	His	Arg	Leu	Ser	Thr	Val
690					695						700				
Glu	His	Ala	His	Leu	Ile	Val	Val	Leu	Asp	Lys	Gly	Arg	Val	Val	Gln
705					710				715						720
Gln	Gly	Thr	His	Gln	Gln	Leu	Leu	Ala	Gln	Gly	Gly	Leu	Tyr	Ala	Lys
			725					730						735	

Leu	Val	Gln	Arg	Gln	Met	Leu	Gly	Leu	Gln	Pro	Ala	Ala	Asp	Phe	Thr
			740					745					750		
Ala	Gly	His	Asn	Glu	Pro	Val	Ala	Asn	Gly	Ser	His	Lys	Ala		
		755					760					765			

<210> 7
 <211> 1280
 <212> PRT
 <213> Homo sapiens

<400> 7

Met	Asp	Leu	Glu	Gly	Asp	Arg	Asn	Gly	Gly	Ala	Lys	Lys	Lys	Asn	Phe
1				5					10					15	
Phe	Lys	Leu	Asn	Asn	Lys	Ser	Glu	Lys	Asp	Lys	Lys	Glu	Lys	Lys	Pro
			20					25					30		
Thr	Val	Ser	Val	Phe	Ser	Met	Phe	Arg	Tyr	Ser	Asn	Trp	Leu	Asp	Lys
		35					40					45			
Leu	Tyr	Met	Val	Val	Gly	Thr	Leu	Ala	Ala	Ile	Ile	His	Gly	Ala	Gly
	50					55					60				
Leu	Pro	Leu	Met	Met	Leu	Val	Phe	Gly	Glu	Met	Thr	Asp	Ile	Phe	Ala
65					70					75					80
Asn	Ala	Gly	Asn	Leu	Glu	Asp	Leu	Met	Ser	Asn	Ile	Thr	Asn	Arg	Ser
				85					90					95	
Asp	Ile	Asn	Asp	Thr	Gly	Phe	Phe	Met	Asn	Leu	Glu	Glu	Asp	Met	Thr
		100						105					110		
Arg	Tyr	Ala	Tyr	Tyr	Tyr	Ser	Gly	Ile	Gly	Ala	Gly	Val	Leu	Val	Ala
	115						120						125		
Ala	Tyr	Ile	Gln	Val	Ser	Phe	Trp	Cys	Leu	Ala	Ala	Gly	Arg	Gln	Ile
	130					135					140				
His	Lys	Ile	Arg	Lys	Gln	Phe	Phe	His	Ala	Ile	Met	Arg	Gln	Glu	Ile
145					150					155					160
Gly	Trp	Phe	Asp	Val	His	Asp	Val	Gly	Glu	Leu	Asn	Thr	Arg	Leu	Thr
			165						170					175	
Asp	Asp	Val	Ser	Lys	Ile	Asn	Glu	Gly	Ile	Gly	Asp	Lys	Ile	Gly	Met
		180						185					190		
Phe	Phe	Gln	Ser	Met	Ala	Thr	Phe	Phe	Thr	Gly	Phe	Ile	Val	Gly	Phe
	195						200					205			
Thr	Arg	Gly	Trp	Lys	Leu	Thr	Leu	Val	Ile	Leu	Ala	Ile	Ser	Pro	Val
	210					215					220				
Leu	Gly	Leu	Ser	Ala	Ala	Val	Trp	Ala	Lys	Ile	Leu	Ser	Ser	Phe	Thr
225					230					235					240
Asp	Lys	Glu	Leu	Leu	Ala	Tyr	Ala	Lys	Ala	Gly	Ala	Val	Ala	Glu	Glu
				245					250					255	
Val	Leu	Ala	Ala	Ile	Arg	Thr	Val	Ile	Ala	Phe	Gly	Gly	Gln	Lys	Lys
		260						265					270		
Glu	Leu	Glu	Arg	Tyr	Asn	Lys	Asn	Leu	Glu	Glu	Ala	Lys	Arg	Ile	Gly
	275						280					285			
Ile	Lys	Lys	Ala	Ile	Thr	Ala	Asn	Ile	Ser	Ile	Gly	Ala	Ala	Phe	Leu
	290					295					300				
Leu	Ile	Tyr	Ala	Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Thr	Leu
305					310					315					320
Val	Leu	Ser	Gly	Glu	Tyr	Ser	Ile	Gly	Gln	Val	Leu	Thr	Val	Phe	Phe
				325					330					335	

Ser Val Leu Ile Gly Ala Phe Ser Val Gly Gln Ala Ser Pro Ser Ile
 340 345 350
 Glu Ala Phe Ala Asn Ala Arg Gly Ala Ala Tyr Glu Ile Phe Lys Ile
 355 360 365
 Ile Asp Asn Lys Pro Ser Ile Asp Ser Tyr Ser Lys Ser Gly His Lys
 370 375 380
 Pro Asp Asn Ile Lys Gly Asn Leu Glu Phe Arg Asn Val His Phe Ser
 385 390 395 400
 Tyr Pro Ser Arg Lys Glu Val Lys Ile Leu Lys Gly Leu Asn Leu Lys
 405 410 415
 Val Gln Ser Gly Gln Thr Val Ala Leu Val Gly Asn Ser Gly Cys Gly
 420 425 430
 Lys Ser Thr Thr Val Gln Leu Met Gln Arg Leu Tyr Asp Pro Thr Glu
 435 440 445
 Gly Met Val Ser Val Asp Gly Gln Asp Ile Arg Thr Ile Asn Val Arg
 450 455 460
 Phe Leu Arg Glu Ile Ile Gly Val Val Ser Gln Glu Pro Val Leu Phe
 465 470 475 480
 Ala Thr Thr Ile Ala Glu Asn Ile Arg Tyr Gly Arg Glu Asn Val Thr
 485 490 495
 Met Asp Glu Ile Glu Lys Ala Val Lys Glu Ala Asn Ala Tyr Asp Phe
 500 505 510
 Ile Met Lys Leu Pro His Lys Phe Asp Thr Leu Val Gly Glu Arg Gly
 515 520 525
 Ala Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala
 530 535 540
 Leu Val Arg Asn Pro Lys Ile Leu Leu Leu Asp Glu Ala Thr Ser Ala
 545 550 555 560
 Leu Asp Thr Glu Ser Glu Ala Val Val Gln Val Ala Leu Asp Lys Ala
 565 570 575
 Arg Lys Gly Arg Thr Thr Ile Val Ile Ala His Arg Leu Ser Thr Val
 580 585 590
 Arg Asn Ala Asp Val Ile Ala Gly Phe Asp Asp Gly Val Ile Val Glu
 595 600 605
 Lys Gly Asn His Asp Glu Leu Met Lys Glu Lys Gly Ile Tyr Phe Lys
 610 615 620
 Leu Val Thr Met Gln Thr Ala Gly Asn Glu Val Glu Leu Glu Asn Ala
 625 630 635 640
 Ala Asp Glu Ser Lys Ser Glu Ile Asp Ala Leu Glu Met Ser Ser Asn
 645 650 655
 Asp Ser Arg Ser Ser Leu Ile Arg Lys Arg Ser Thr Arg Arg Ser Val
 660 665 670
 Arg Gly Ser Gln Ala Gln Asp Arg Lys Leu Ser Thr Lys Glu Ala Leu
 675 680 685
 Asp Glu Ser Ile Pro Pro Val Ser Phe Trp Arg Ile Met Lys Leu Asn
 690 695 700
 Leu Thr Glu Trp Pro Tyr Phe Val Val Gly Val Phe Cys Ala Ile Ile
 705 710 715 720
 Asn Gly Gly Leu Gln Pro Ala Phe Ala Ile Ile Phe Ser Lys Ile Ile
 725 730 735
 Gly Val Phe Thr Arg Ile Asp Asp Pro Glu Thr Lys Arg Gln Asn Ser
 740 745 750
 Asn Leu Phe Ser Leu Leu Phe Leu Ala Leu Gly Ile Ile Ser Phe Ile
 755 760 765

Thr Phe Phe Leu Gln Gly Phe Thr Phe Gly Lys Ala Gly Glu Ile Leu
 770 775 780
 Thr Lys Arg Leu Arg Tyr Met Val Phe Arg Ser Met Leu Arg Gln Asp
 785 790 795 800
 Val Ser Trp Phe Asp Asp Pro Lys Asn Thr Thr Gly Ala Leu Thr Thr
 805 810 815
 Arg Leu Ala Asn Asp Ala Ala Gln Val Lys Gly Ala Ile Gly Ser Arg
 820 825 830
 Leu Ala Val Ile Thr Gln Asn Ile Ala Asn Leu Gly Thr Gly Ile Ile
 835 840 845
 Ile Ser Phe Ile Tyr Gly Trp Gln Leu Thr Leu Leu Leu Ala Ile
 850 855 860
 Val Pro Ile Ile Ala Ile Ala Gly Val Val Glu Met Lys Met Leu Ser
 865 870 875 880
 Gly Gln Ala Leu Lys Asp Lys Lys Glu Leu Glu Gly Ala Gly Lys Ile
 885 890 895
 Ala Thr Glu Ala Ile Glu Asn Phe Arg Thr Val Val Ser Leu Thr Gln
 900 905 910
 Glu Gln Lys Phe Glu His Met Tyr Ala Gln Ser Leu Gln Val Pro Tyr
 915 920 925
 Arg Asn Ser Leu Arg Lys Ala His Ile Phe Gly Ile Thr Phe Ser Phe
 930 935 940
 Thr Gln Ala Met Met Tyr Phe Ser Tyr Ala Gly Cys Phe Arg Phe Gly
 945 950 955 960
 Ala Tyr Leu Val Ala His Lys Leu Met Ser Phe Glu Asp Val Leu Leu
 965 970 975
 Val Phe Ser Ala Val Val Phe Gly Ala Met Ala Val Gly Gln Val Ser
 980 985 990
 Ser Phe Ala Pro Asp Tyr Ala Lys Ala Lys Ile Ser Ala Ala His Ile
 995 1000 1005
 Ile Met Ile Ile Glu Lys Thr Pro Leu Ile Asp Ser Tyr Ser Thr Glu
 1010 1015 1020
 Gly Leu Met Pro Asn Thr Leu Glu Gly Asn Val Thr Phe Gly Glu Val
 1025 1030 1035 1040
 Val Phe Asn Tyr Pro Thr Arg Pro Asp Ile Pro Val Leu Gln Gly Leu
 1045 1050 1055
 Ser Leu Glu Val Lys Lys Gly Gln Thr Leu Ala Leu Val Gly Ser Ser
 1060 1065 1070
 Gly Cys Gly Lys Ser Thr Val Val Gln Leu Leu Glu Arg Phe Tyr Asp
 1075 1080 1085
 Pro Leu Ala Gly Lys Val Leu Leu Asp Gly Lys Glu Ile Lys Arg Leu
 1090 1095 1100
 Asn Val Gln Trp Leu Arg Ala His Leu Gly Ile Val Ser Gln Glu Pro
 1105 1110 1115 1120
 Ile Leu Phe Asp Cys Ser Ile Ala Glu Asn Ile Ala Tyr Gly Asp Asn
 1125 1130 1135
 Ser Arg Val Val Ser Gln Glu Glu Ile Val Arg Ala Ala Lys Glu Ala
 1140 1145 1150
 Asn Ile His Ala Phe Ile Glu Ser Leu Pro Asn Lys Tyr Ser Thr Lys
 1155 1160 1165
 Val Gly Asp Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile
 1170 1175 1180
 Ala Ile Ala Arg Ala Leu Val Arg Gln Pro His Ile Leu Leu Leu Asp
 1185 1190 1195 1200

Glu	Ala	Thr	Ser	Ala	Leu	Asp	Thr	Glu	Ser	Glu	Lys	Val	Val	Gln	Glu
				1205					1210					1215	
Ala	Leu	Asp	Lys	Ala	Arg	Glu	Gly	Arg	Thr	Cys	Ile	Val	Ile	Ala	His
			1220					1225						1230	
Arg	Leu	Ser	Thr	Ile	Gln	Asn	Ala	Asp	Leu	Ile	Val	Val	Phe	Gln	Asn
			1235					1240						1245	
Gly	Arg	Val	Lys	Glu	His	Gly	Thr	His	Gln	Gln	Leu	Leu	Ala	Gln	Lys
			1250					1255						1260	
Gly	Ile	Tyr	Phe	Ser	Met	Val	Ser	Val	Gln	Ala	Gly	Thr	Lys	Arg	Gln
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<210> 8

<211> 2001

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> 30, 70, 280, 477, 558, 1471, 1651, 1689, 1724

<223> Xaa = Any Amino Acid

<221> VARIANT

<222> 30, 70, 280, 477, 558, 1471, 1651, 1689, 1724

<223> Xaa = Any Amino Acid

<400> 8

Met	Ser	Ser	Leu	Gly	Phe	Thr	Ser	Lys	Glu	Gln	Arg	Asn	Leu	Gly	Leu
1				5					10					15	
Leu	Val	His	Leu	Met	Thr	Ser	Asn	Pro	Lys	Ile	Leu	Tyr	Xaa	Pro	Ala
			20					25					30		
Gly	Ser	Glu	Val	Asp	Arg	Val	Ile	Leu	Lys	Ala	Asn	Glu	Thr	Phe	Ala
			35				40					45			
Phe	Val	Gly	Asn	Val	Thr	His	Tyr	Ala	Gln	Val	Trp	Leu	Asn	Ile	Ser
			50			55					60				
Ala	Glu	Ile	Arg	Ser	Xaa	Leu	Glu	Gln	Gly	Arg	Leu	Gln	Gln	His	Leu
65					70				75					80	
Arg	Trp	Leu	Gln	Gln	Tyr	Val	Ala	Glu	Leu	Arg	Pro	His	Pro	Glu	Ala
			85					90					95		
Leu	Asn	Leu	Ser	Leu	Asp	Glu	Leu	Pro	Pro	Ala	Leu	Arg	Gln	Asp	Asn
			100					105					110		
Phe	Ser	Leu	Pro	Ser	Gly	Met	Ala	Leu	Leu	Gln	Gln	Leu	Asp	Thr	Ile
			115				120						125		
Asp	Asn	Ala	Pro	Cys	Gly	Trp	Ile	Gln	Phe	Met	Ser	Lys	Val	Ser	Val
			130				135					140			
Asp	Ile	Phe	Lys	Gly	Phe	Pro	Asp	Glu	Glu	Ser	Ile	Val	Asn	Tyr	Thr
145					150					155				160	
Leu	Asn	Gln	Ala	Tyr	Gln	Asp	Asn	Val	Thr	Val	Phe	Ala	Gly	Val	Ile
				165					170					175	
Phe	Gln	Thr	Arg	Lys	Asp	Gly	Ser	Leu	Pro	Pro	His	Val	His	Tyr	Lys
			180					185					190		
Ile	Arg	Gln	Asn	Ser	Ser	Phe	Thr	Glu	Lys	Thr	Asn	Glu	Ile	Arg	Arg
			195				200					205			
Ala	Tyr	Trp	Arg	Pro	Gly	Pro	Asn	Thr	Gly	Gly	Arg	Phe	Tyr	Phe	Leu

210		215		220
Tyr Gly Phe Val Trp	Ile Gln Asp Met Met	Glu Arg Ala Ile Ile Asp		
225	230	235	240	
Thr Phe Val Gly His Asp	Val Val Glu Pro Gly	Ser Tyr Val Gln Met		
	245	250	255	
Phe Pro Tyr Pro Cys Tyr	Thr Arg Asp Asp	Phe Leu Phe Val Ile Glu		
	260	265	270	
His Met Met Pro Leu Cys	Met Xaa Ile Ser Trp	Val Tyr Ser Val Ala		
	275	280	285	
Met Thr Ile Gln His Ile	Val Ala Glu Lys Glu	His Arg Leu Lys Glu		
	290	295	300	
Val Met Lys Thr Met Gly	Leu Asn Asn Ala Val	His Trp Val Ala Trp		
305	310	315	320	
Phe Ile Thr Gly Phe Val	Gln Leu Ser Ile Ser	Val Thr Ala Leu Thr		
	325	330	335	
Ala Ile Leu Lys Tyr Gly	Gln Val Leu Met His	Ser His Val Val Ile		
	340	345	350	
Ile Trp Leu Phe Leu Ala	Val Tyr Ala Val Ala	Thr Ile Met Phe Cys		
	355	360	365	
Phe Leu Val Ser Val Leu	Tyr Ser Lys Ala Lys	Leu Ala Ser Ala Gly		
	370	375	380	
Gly Ile Ile Tyr Phe Leu	Ser Tyr Val Pro Tyr	Met Tyr Val Ala Ile		
385	390	395	400	
Arg Glu Glu Val Ala His	Asp Lys Ile Thr Ala	Phe Glu Lys Cys Ile		
	405	410	415	
Ala Ser Leu Met Ser Thr	Thr Ala Phe Gly Leu	Gly Ser Lys Tyr Phe		
	420	425	430	
Ala Leu Tyr Glu Val Ala	Gly Val Gly Ile Gln	Trp His Thr Phe Ser		
	435	440	445	
Gln Ser Pro Val Glu Gly	Asp Phe Asn Leu Leu	Leu Ala Val Thr		
	450	455	460	
Met Leu Met Val Asp Ala	Val Val Tyr Gly Ile	Leu Xaa Trp Tyr Ile		
465	470	475	480	
Glu Ala Val His Pro Gly	Met Tyr Gly Leu Pro	Arg Pro Trp Tyr Phe		
	485	490	495	
Pro Leu Gln Lys Ser Tyr	Trp Leu Gly Ser Gly	Arg Thr Glu Ala Trp		
	500	505	510	
Glu Trp Ser Trp Pro Trp	Ala Arg Thr Pro Arg	Leu Ser Val Met Glu		
	515	520	525	
Glu Asp Gln Ala Cys Ala	Met Glu Ser Arg Arg	Phe Glu Glu Thr Arg		
	530	535	540	
Gly Met Glu Glu Glu Pro	Thr His Leu Pro Leu	Val Val Xaa Val Asp		
545	550	555	560	
Lys Leu Thr Lys Val Tyr	Lys Asp Asp Lys Lys	Leu Ala Leu Asn Lys		
	565	570	575	
Leu Ser Leu Asn Leu Tyr	Glu Asn Gln Gly Val	Ser Phe Leu Gly His		
	580	585	590	
Asn Gly Ala Gly Lys Thr	Thr Thr Met Ser Ile	Leu Thr Gly Leu Phe		
	595	600	605	
Pro Pro Thr Ser Gly Ser	Ala Thr Ile Tyr Gly	His Asp Ile Arg Thr		
	610	615	620	
Glu Met Asp Glu Ile Arg	Lys Asn Gly His Val	Pro Gln His Asn Val		
625	630	635	640	
Leu Phe Asp Arg Leu Thr	Val Glu Glu His Leu	Trp Phe Tyr Ser Arg		

				645					650				655			
Leu	Lys	Ser	Met	Ala	Gln	Glu	Glu	Ile	Pro	Arg	Glu	Met	Asp	Lys	Met	
			660					665					670			
Ile	Glu	Asp	Leu	Glu	Leu	Ser	Asn	Lys	Arg	His	Ser	Leu	Val	Gln	Thr	
		675					680					685				
Leu	Ser	Gly	Gly	Met	Lys	Arg	Lys	Val	Ser	Val	Ala	Ile	Ala	Phe	Val	
	690					695					700					
Gly	Gly	Ser	Arg	Ala	Ile	Ile	Leu	Asp	Glu	Pro	Thr	Ala	Gly	Val	Asp	
705					710				715						720	
Pro	Tyr	Ala	Arg	Arg	Ala	Ile	Trp	Asp	Leu	Ile	Leu	Lys	Tyr	Lys	Pro	
			725					730							735	
Gly	Arg	Thr	Ile	Leu	Leu	Ser	Thr	His	His	Met	Asp	Glu	Ala	Asp	Leu	
		740						745					750			
Leu	Gly	Asp	Arg	Ile	Ala	Ile	Ile	Ser	His	Gly	Lys	Leu	Lys	Cys	Cys	
	755						760					765				
Gly	Ser	Pro	Leu	Phe	Leu	Lys	Gly	Thr	Tyr	Gly	Asp	Gly	Tyr	Arg	Leu	
	770					775					780					
Thr	Leu	Val	Lys	Arg	Pro	Ala	Glu	Pro	Gly	Gly	Pro	Gln	Glu	Pro	Gly	
785					790						795				800	
Leu	Ala	Ser	Ser	Pro	Pro	Gly	Arg	Ala	Pro	Leu	Ser	Ser	Cys	Ser	Glu	
			805					810						815		
Leu	Gln	Val	Ser	Gln	Phe	Ile	Arg	Lys	His	Val	Ala	Ser	Cys	Leu	Leu	
		820						825					830			
Val	Ser	Asp	Thr	Ser	Thr	Glu	Leu	Ser	Tyr	Ile	Leu	Pro	Ser	Glu	Ala	
	835						840					845				
Ala	Lys	Lys	Gly	Ala	Phe	Glu	Arg	Leu	Phe	Gln	His	Leu	Glu	Arg	Ser	
	850					855					860					
Leu	Asp	Ala	Leu	His	Leu	Ser	Ser	Phe	Gly	Leu	Met	Asp	Thr	Thr	Leu	
865					870					875					880	
Glu	Glu	Val	Phe	Leu	Lys	Val	Ser	Gly	Gly	Asp	Gln	Ser	Leu	Glu	Asn	
			885					890						895		
Ser	Gly	Ala	Asp	Val	Lys	Glu	Ser	Arg	Lys	Asp	Val	Leu	Pro	Gly	Ala	
		900						905					910			
Glu	Gly	His	Ala	Ser	Gly	Glu	Gly	His	Ala	Gly	Asn	Leu	Ala	Arg	Cys	
	915						920					925				
Ser	Glu	Leu	Thr	Gln	Ser	Gln	Ala	Ser	Leu	Gln	Ser	Ala	Ser	Ser	Val	
	930					935					940					
Gly	Ser	Ala	Leu	Gly	Asp	Glu	Gly	Ala	Gly	Tyr	Thr	Asp	Val	Tyr	Gly	
945					950					955					960	
Asp	Tyr	Pro	Pro	Leu	Phe	Asp	Asn	Pro	Gln	Asp	Pro	Asp	Asn	Val	Ser	
			965					970						975		
Leu	Gln	Glu	Val	Glu	Ala	Glu	Ala	Leu	Ser	Arg	Val	Gly	Gln	Gly	Ser	
		980						985					990			
Arg	Lys	Leu	Asp	Gly	Gly	Trp	Leu	Lys	Val	Arg	Gln	Phe	His	Gly	Leu	
	995					1000						1005				
Leu	Val	Lys	Arg	Phe	His	Cys	Ala	Arg	Arg	Asn	Ser	Lys	Ala	Leu	Phe	
	1010					1015					1020					
Ser	Gln	Ile	Leu	Leu	Pro	Ala	Phe	Phe	Val	Cys	Val	Ala	Met	Thr	Val	
1025					1030					1035					1040	
Ala	Leu	Ser	Val	Pro	Glu	Ile	Gly	Asp	Leu	Pro	Pro	Leu	Val	Leu	Ser	
			1045					1050					1055			
Pro	Ser	Gln	Tyr	His	Asn	Tyr	Thr	Gln	Pro	Arg	Gly	Asn	Phe	Ile	Pro	
		1060						1065				1070				
Tyr	Ala	Asn	Glu	Glu	Arg	Arg	Glu	Tyr	Arg	Leu	Arg	Leu	Ser	Pro	Asp	

1075	1080	1085
Ala Ser Pro Gln Gln Leu Val Ser Thr Phe Arg Leu Pro Ser Gly Val		
1090	1095	1100
Gly Ala Thr Cys Val Leu Lys Ser Pro Ala Asn Gly Ser Leu Gly Pro		
1105	1110	1115
Thr Leu Asn Leu Ser Ser Gly Glu Ser Arg Leu Leu Ala Ala Arg Phe		
1125	1130	1135
Phe Asp Ser Met Cys Leu Glu Ser Phe Thr Gln Gly Leu Pro Leu Ser		
1140	1145	1150
Asn Phe Val Pro Pro Pro Pro Ser Pro Ala Pro Ser Asp Ser Pro Ala		
1155	1160	1165
Ser Pro Asp Glu Asp Leu Gln Ala Trp Asn Val Ser Leu Pro Pro Thr		
1170	1175	1180
Ala Gly Gln Glu Met Trp Thr Ser Ala Pro Ser Leu Pro Arg Leu Val		
1185	1190	1195
Arg Glu Pro Val Arg Cys Thr Cys Ser Ala Gln Gly Thr Gly Phe Ser		
1205	1210	1215
Cys Pro Asn Ser Val Gly Gly His Pro Pro Gln Met Arg Val Val Thr		
1220	1225	1230
Gly Asp Ile Leu Thr Asp Ile Thr Gly His Asn Val Ser Glu Tyr Leu		
1235	1240	1245
Leu Phe Thr Ser Asp Arg Phe Arg Leu His Arg Tyr Gly Ala Ile Thr		
1250	1255	1260
Phe Gly Asn Val Leu Lys Ser Ile Pro Ala Ser Phe Gly Thr Arg Ala		
1265	1270	1275
Pro Pro Met Val Arg Lys Ile Arg Cys Ala Arg Ala Ala Gln Val Phe		
1285	1290	1295
Tyr Asn Asn Lys Gly Tyr His Ser Met Pro Thr Tyr Leu Asn Ser Leu		
1300	1305	1310
Asn Asn Ala Ile Leu Arg Ala Asn Leu Pro Lys Ser Lys Gly Asn Pro		
1315	1320	1325
Ala Ala Tyr Gly Ile Thr Val Thr Asn His Pro Met Asn Lys Thr Ser		
1330	1335	1340
Ala Ser Leu Ser Leu Asp Tyr Leu Leu Gln Gly Thr Asp Val Val Ile		
1345	1350	1355
Ala Ile Phe Ile Ile Val Ala Met Ser Phe Val Pro Ala Ser Phe Val		
1365	1370	1375
Val Phe Leu Val Ala Glu Lys Ser Thr Lys Ala Lys His Leu Gln Phe		
1380	1385	1390
Val Ser Gly Cys Asn Pro Ile Ile Tyr Trp Leu Ala Asn Tyr Val Trp		
1395	1400	1405
Asp Met Leu Asn Tyr Leu Val Pro Ala Thr Cys Cys Val Ile Ile Leu		
1410	1415	1420
Phe Val Phe Asp Leu Pro Ala Tyr Thr Ser Pro Thr Asn Phe Pro Ala		
1425	1430	1435
Val Leu Ser Leu Phe Leu Leu Tyr Gly Trp Ser Ile Thr Pro Ile Met		
1445	1450	1455
Tyr Pro Ala Ser Phe Trp Phe Glu Val Pro Ser Ser Ala Tyr Xaa Phe		
1460	1465	1470
Leu Ile Val Ile Asn Leu Phe Ile Gly Ile Thr Ala Thr Val Ala Thr		
1475	1480	1485
Phe Leu Leu Gln Leu Phe Glu His Asp Lys Asp Leu Lys Val Val Asn		
1490	1495	1500
Ser Tyr Leu Lys Ser Cys Phe Leu Ile Phe Pro Asn Tyr Asn Leu Gly		

1505		1510		1515		1520
His Gly Leu Met	Glu Met Ala Tyr Asn	Glu Tyr Ile Asn	Glu Tyr Tyr			
	1525		1530		1535	
Ala Lys Ile Gly	Gln Phe Asp Lys Met	Lys Ser Pro Phe	Glu Trp Asp			
	1540		1545		1550	
Ile Val Thr Arg	Gly Leu Val Ala Met	Ala Val Glu Gly	Val Val Gly			
	1555		1560		1565	
Phe Leu Leu Thr	Ile Met Cys Gln Tyr	Asn Phe Leu Arg	Arg Pro Gln			
	1570		1575		1580	
Arg Met Pro Val	Ser Thr Lys Pro Val	Glu Asp Asp Val	Asp Val Ala			
1585		1590		1595		1600
Ser Glu Arg Gln	Arg Val Leu Arg	Gly Asp Ala Asp	Asn Asp Met Val			
	1605		1610		1615	
Lys Ile Glu Asn	Leu Thr Lys Val Tyr	Lys Ser Arg Lys	Ile Gly Arg			
	1620		1625		1630	
Ile Leu Ala Val	Asp Arg Leu Cys Leu	Gly Val Arg Pro	Gly Glu Cys			
	1635		1640		1645	
Phe Gly Xaa Leu	Gly Val Asn Gly Ala	Gly Lys Thr Ser	Thr Phe Lys			
	1650		1655		1660	
Met Leu Thr Gly	Asp Glu Ser Thr Thr	Gly Gly Glu Ala	Phe Val Asn			
1665		1670		1675		1680
Gly His Ser Val	Leu Lys Glu Leu Xaa	Gln Val Gln Gln	Ser Leu Gly			
	1685		1690		1695	
Tyr Cys Pro Gln	Cys Asp Ala Leu Phe	Asp Glu Leu Thr	Ala Arg Glu			
	1700		1705		1710	
His Leu Gln Leu	Tyr Thr Arg Leu Arg	Gly Ile Xaa Trp	Lys Asp Glu			
	1715		1720		1725	
Ala Arg Val Val	Lys Trp Ala Leu Glu	Lys Leu Glu Leu	Thr Lys Tyr			
	1730		1735		1740	
Ala Asp Lys Pro	Ala Gly Thr Tyr Ser	Gly Gly Asn Lys	Arg Lys Leu			
1745		1750		1755		1760
Ser Thr Ala Ile	Ala Leu Ile Gly Tyr	Pro Ala Phe Ile	Phe Leu Asp			
	1765		1770		1775	
Glu Pro Thr Thr	Gly Met Asp Pro Lys	Ala Arg Arg Phe	Leu Trp Asn			
	1780		1785		1790	
Leu Ile Leu Asp	Leu Ile Lys Thr Gly	Arg Ser Val Val	Leu Thr Ser			
	1795		1800		1805	
His Ser Met Glu	Glu Cys Glu Ala Leu	Cys Thr Arg Leu	Ala Ile Met			
	1810		1815		1820	
Val Asn Gly Arg	Leu Arg Cys Leu Gly	Ser Ile Gln His	Leu Lys Asn			
1825		1830		1835		1840
Arg Phe Gly Asp	Gly Tyr Met Ile Thr	Val Arg Thr Lys	Ser Ser Gln			
	1845		1850		1855	
Ser Val Lys Asp	Val Val Arg Phe Phe	Asn Arg Asn Phe	Pro Glu Ala			
	1860		1865		1870	
Met Leu Lys Glu	Arg His His Thr Lys	Val Gln Tyr Gln	Leu Lys Ser			
	1875		1880		1885	
Glu His Ile Ser	Leu Ala Gln Val Phe	Ser Lys Met Glu	Gln Val Ser			
	1890		1895		1900	
Gly Val Leu Gly	Ile Glu Asp Tyr Ser	Val Ser Gln Thr	Thr Leu Asp			
1905		1910		1915		1920
Asn Val Phe Val	Asn Phe Ala Lys Lys	Gln Ser Asp Asn	Leu Glu Gln			
	1925		1930		1935	
Gln Glu Thr Glu	Pro Pro Ser Ala Leu	Gln Ser Pro Leu	Gly Cys Leu			

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<210> 9
<211> 646
<212> PRT
<213> Homo sapiens
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Met	Ala	Glu	Lys	Ala	Leu	Glu	Ala	Val	Gly	Cys	Gly	Leu	Gly	Pro	Gly	
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Ala	Val	Ala	Met	Ala	Val	Thr	Leu	Glu	Asp	Gly	Ala	Glu	Pro	Pro	Val	
			20					25					30			
Leu	Thr	Thr	His	Leu	Lys	Lys	Val	Glu	Asn	His	Ile	Thr	Glu	Ala	Gln	
			35				40					45				
Arg	Phe	Ser	His	Leu	Pro	Lys	Arg	Ser	Ala	Val	Asp	Ile	Glu	Phe	Val	
	50					55					60					
Glu	Leu	Ser	Tyr	Ser	Val	Arg	Glu	Gly	Pro	Cys	Trp	Arg	Lys	Arg	Gly	
65					70					75					80	
Tyr	Lys	Thr	Leu	Leu	Lys	Cys	Leu	Ser	Gly	Lys	Phe	Cys	Arg	Arg	Glu	
				85					90					95		
Leu	Ile	Gly	Ile	Met	Gly	Pro	Ser	Gly	Ala	Gly	Lys	Ser	Thr	Phe	Met	
			100					105					110			
Asn	Ile	Leu	Ala	Gly	Tyr	Arg	Glu	Ser	Gly	Met	Lys	Gly	Gln	Ile	Leu	
		115					120					125				
Val	Asn	Gly	Arg	Pro	Arg	Glu	Leu	Arg	Thr	Phe	Arg	Lys	Met	Ser	Cys	
	130					135					140					
Tyr	Ile	Met	Gln	Asp	Asp	Met	Leu	Leu	Pro	His	Leu	Thr	Val	Leu	Glu	
145					150					155					160	
Ala	Met	Met	Val	Ser	Ala	Asn	Leu	Asn	Leu	Thr	Glu	Asn	Pro	Asp	Val	
				165					170					175		
Lys	Asn	Asp	Leu	Val	Thr	Glu	Ile	Leu	Thr	Ala	Leu	Gly	Leu	Met	Ser	
			180					185					190			
Cys	Ser	His	Thr	Arg	Thr	Ala	Leu	Leu	Ser	Gly	Gly	Gln	Arg	Lys	Arg	
		195					200					205				
Leu	Ala	Ile	Ala	Leu	Glu	Leu	Val	Asn	Asn	Pro	Pro	Val	Met	Phe	Phe	
	210					215					220					
Asp	Glu	Pro	Thr	Ser	Gly	Leu	Asp	Ser	Ala	Ser	Cys	Phe	Gln	Val	Val	
225					230					235					240	
Ser	Leu	Met	Lys	Ser	Leu	Ala	Gln	Gly	Gly	Arg	Thr	Ile	Ile	Cys	Thr	
				245					250					255		
Ile	His	Gln	Pro	Ser	Ala	Lys	Leu	Phe	Glu	Met	Phe	Asp	Lys	Leu	Tyr	
			260					265					270			
Ile	Leu	Ser	Gln	Gly	Gln	Cys	Ile	Phe	Lys	Gly	Val	Val	Thr	Asn	Leu	
		275					280					285				
Ile	Pro	Tyr	Leu	Lys	Gly	Leu	Gly	Leu	His	Cys	Pro	Thr	Tyr	His	Asn	

290	295	300
Pro Ala Asp Phe Ile Ile Glu Val Ala Ser Gly Glu Tyr Gly Asp Leu		
305	310	315
Asn Pro Met Leu Phe Arg Ala Val Gln Asn Gly Leu Cys Ala Met Ala		320
	325	330
Glu Lys Lys Ser Ser Pro Glu Lys Asn Glu Val Pro Ala Pro Cys Pro		335
	340	345
Pro Cys Pro Pro Glu Val Asp Pro Ile Glu Ser His Thr Phe Ala Thr		350
	355	360
Ser Thr Leu Thr Gln Phe Cys Ile Leu Phe Lys Arg Thr Phe Leu Ser		365
	370	375
Ile Leu Arg Asp Thr Val Leu Thr His Leu Arg Phe Met Ser His Val		380
385	390	395
Val Ile Gly Val Leu Ile Gly Leu Leu Tyr Leu His Ile Gly Asp Asp		400
	405	410
Ala Ser Lys Val Phe Asn Asn Thr Gly Cys Leu Phe Phe Ser Met Leu		415
	420	425
Phe Leu Met Phe Ala Ala Leu Met Pro Thr Val Leu Thr Phe Pro Leu		430
	435	440
Glu Met Ala Val Phe Met Arg Glu His Leu Asn Tyr Trp Tyr Ser Leu		445
	450	455
Lys Ala Tyr Tyr Leu Ala Lys Thr Met Ala Asp Val Pro Phe Gln Val		460
465	470	475
Val Cys Pro Val Val Tyr Cys Ser Ile Val Tyr Trp Met Thr Gly Gln		480
	485	490
Pro Ala Glu Thr Ser Arg Phe Leu Leu Phe Ser Ala Leu Ala Thr Ala		495
	500	505
Thr Ala Leu Val Ala Gln Ser Leu Gly Leu Leu Ile Gly Ala Ala Ser		510
	515	520
Asn Ser Leu Gln Val Ala Thr Phe Val Gly Pro Val Thr Ala Ile Pro		525
	530	535
Val Leu Leu Phe Ser Gly Phe Phe Val Ser Phe Lys Thr Ile Pro Thr		540
545	550	555
Tyr Leu Gln Trp Ser Ser Tyr Leu Ser Tyr Val Arg Tyr Gly Phe Glu		560
	565	570
Gly Val Ile Leu Thr Ile Tyr Gly Met Glu Arg Gly Asp Leu Thr Cys		575
	580	585
Leu Glu Glu Arg Cys Pro Phe Arg Glu Pro Gln Ser Ile Leu Arg Ala		590
	595	600
Leu Asp Val Glu Asp Ala Lys Leu Tyr Met Asp Phe Leu Val Leu Gly		605
	610	615
Ile Phe Phe Leu Ala Leu Arg Leu Leu Ala Tyr Leu Val Leu Arg Tyr		620
625	630	635
Arg Val Lys Ser Glu Arg		640
	645	

<210> 10

<211> 638

<212> PRT

<213> Homo sapiens

<400> 10

Met Glu Ala Thr Glu Thr Asp Leu Leu Asn Gly His Leu Lys Lys Val

1	5	10	15
Asp Asn Asn Leu Thr Glu Ala Gln Arg Phe Ser Ser Leu Pro Arg Arg			
20	25	30	
Ala Ala Val Asn Ile Glu Phe Arg Asp Leu Ser Tyr Ser Val Pro Glu			
35	40	45	
Gly Pro Trp Trp Arg Lys Lys Gly Tyr Lys Thr Leu Leu Lys Gly Ile			
50	55	60	
Ser Gly Lys Phe Asn Ser Gly Glu Leu Val Ala Ile Met Gly Pro Ser			
65	70	75	80
Gly Ala Gly Lys Ser Thr Leu Met Asn Ile Leu Ala Gly Tyr Arg Glu			
85	90	95	
Thr Gly Met Lys Gly Ala Val Leu Ile Asn Gly Leu Pro Arg Asp Leu			
100	105	110	
Arg Cys Phe Arg Lys Val Ser Cys Tyr Ile Met Gln Asp Asp Met Leu			
115	120	125	
Leu Pro His Leu Thr Val Gln Glu Ala Met Met Val Ser Ala His Leu			
130	135	140	
Lys Leu Gln Glu Lys Asp Glu Gly Arg Arg Glu Met Val Lys Glu Ile			
145	150	155	160
Leu Thr Ala Leu Gly Leu Leu Ser Cys Ala Asn Thr Arg Thr Gly Ser			
165	170	175	
Leu Ser Gly Gly Gln Arg Lys Arg Leu Ala Ile Ala Leu Glu Leu Val			
180	185	190	
Asn Asn Pro Pro Val Met Phe Phe Asp Glu Pro Thr Ser Gly Leu Asp			
195	200	205	
Ser Ala Ser Cys Phe Gln Val Val Ser Leu Met Lys Gly Leu Ala Gln			
210	215	220	
Gly Gly Arg Ser Ile Ile Cys Thr Ile His Gln Pro Ser Ala Lys Leu			
225	230	235	240
Phe Glu Leu Phe Asp Gln Leu Tyr Val Leu Ser Gln Gly Gln Cys Val			
245	250	255	
Tyr Arg Gly Lys Val Cys Asn Leu Val Pro Tyr Leu Arg Asp Leu Gly			
260	265	270	
Leu Asn Cys Pro Thr Tyr His Asn Pro Ala Asp Phe Val Met Glu Val			
275	280	285	
Ala Ser Gly Glu Tyr Gly Asp Gln Asn Ser Arg Leu Val Arg Ala Val			
290	295	300	
Arg Glu Gly Met Cys Asp Ser Asp His Lys Arg Asp Leu Gly Gly Asp			
305	310	315	320
Ala Glu Val Asn Pro Phe Leu Trp His Arg Pro Ser Glu Glu Val Lys			
325	330	335	
Gln Thr Lys Arg Leu Lys Gly Leu Arg Lys Asp Ser Ser Ser Met Glu			
340	345	350	
Gly Cys His Ser Phe Ser Ala Ser Cys Leu Thr Gln Phe Cys Ile Leu			
355	360	365	
Phe Lys Arg Thr Phe Leu Ser Ile Met Arg Asp Ser Val Leu Thr His			
370	375	380	
Leu Arg Ile Thr Ser His Ile Gly Ile Gly Leu Leu Ile Gly Leu Leu			
385	390	395	400
Tyr Leu Gly Ile Gly Asn Glu Ala Lys Lys Val Leu Ser Asn Ser Gly			
405	410	415	
Phe Leu Phe Phe Ser Met Leu Phe Leu Met Phe Ala Ala Leu Met Pro			
420	425	430	
Thr Val Leu Thr Phe Pro Leu Glu Met Gly Val Phe Leu Arg Glu His			

	435		440		445										
Leu	Asn	Tyr	Trp	Tyr	Ser	Leu	Lys	Ala	Tyr	Tyr	Leu	Ala	Lys	Thr	Met
	450						455					460			
Ala	Asp	Val	Pro	Phe	Gln	Ile	Met	Phe	Pro	Val	Ala	Tyr	Cys	Ser	Ile
465						470					475				480
Val	Tyr	Trp	Met	Thr	Ser	Gln	Pro	Ser	Asp	Ala	Val	Ala	Phe	Val	Leu
				485					490						495
Phe	Ala	Ala	Leu	Gly	Thr	Met	Thr	Ser	Leu	Val	Ala	Gln	Ser	Leu	Gly
			500					505					510		
Leu	Leu	Ile	Gly	Ala	Ala	Ser	Thr	Ser	Leu	Gln	Val	Ala	Thr	Phe	Val
		515					520					525			
Gly	Pro	Val	Thr	Ala	Ile	Pro	Val	Leu	Leu	Phe	Ser	Gly	Phe	Phe	Val
	530					535					540				
Ser	Phe	Asp	Thr	Ile	Pro	Thr	Tyr	Leu	Gln	Trp	Met	Ser	Tyr	Ile	Ser
545					550					555					560
Tyr	Val	Arg	Tyr	Gly	Phe	Glu	Gly	Val	Ile	Leu	Ser	Ile	Tyr	Gly	Leu
				565					570						575
Asp	Arg	Glu	Asp	Leu	His	Cys	Asp	Ile	Asp	Glu	Thr	Cys	His	Phe	Gln
			580					585					590		
Lys	Ser	Glu	Ala	Ile	Leu	Arg	Glu	Leu	Asp	Val	Glu	Asn	Ala	Lys	Leu
		595					600					605			
Tyr	Leu	Asp	Phe	Ile	Val	Leu	Gly	Ile	Phe	Phe	Ile	Ser	Leu	Arg	Leu
	610					615					620				
Ile	Ala	Tyr	Phe	Val	Leu	Arg	Tyr	Lys	Ile	Arg	Ala	Glu	Arg		
625					630					635					